

FIGURE 1

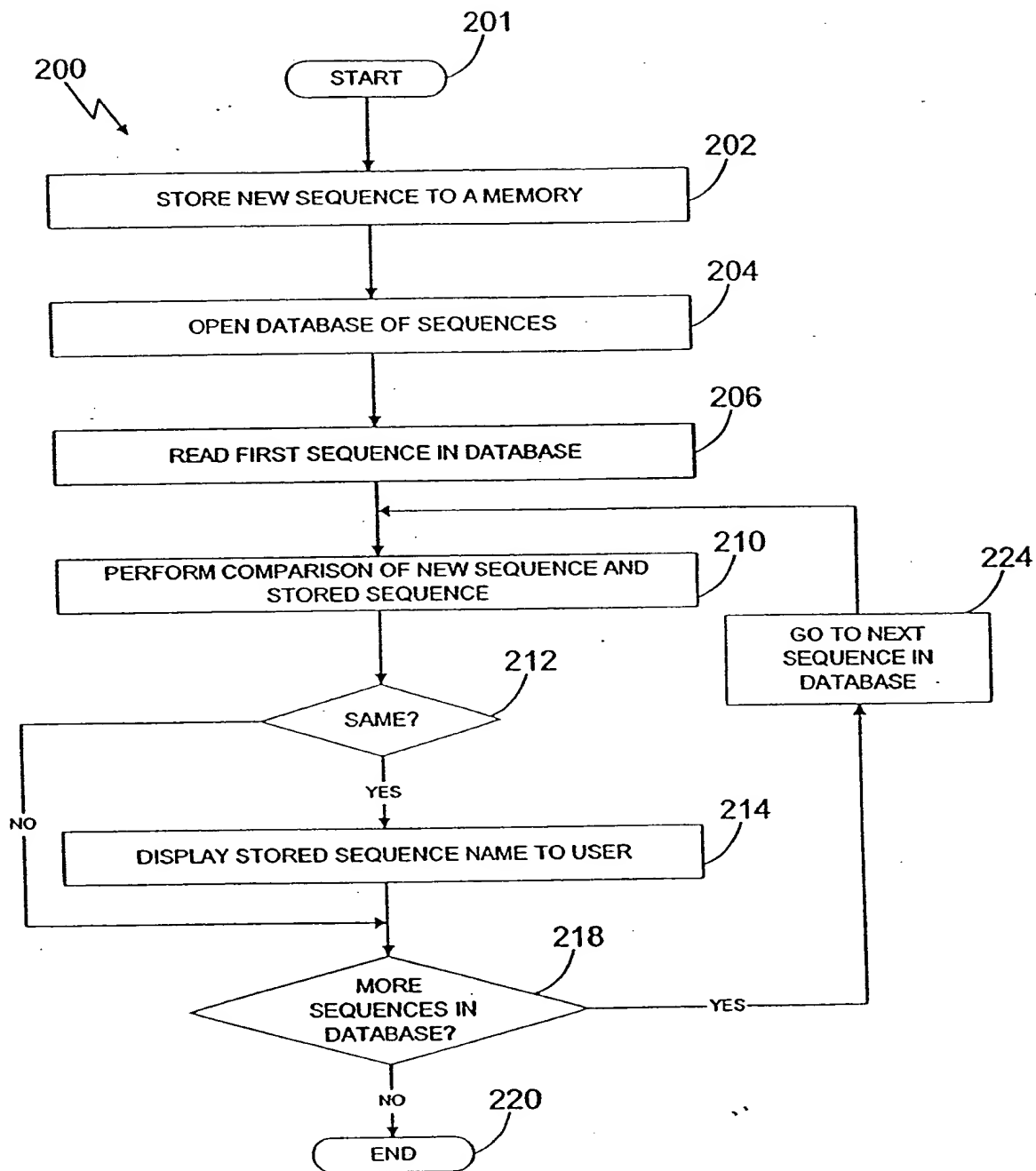


FIGURE 2

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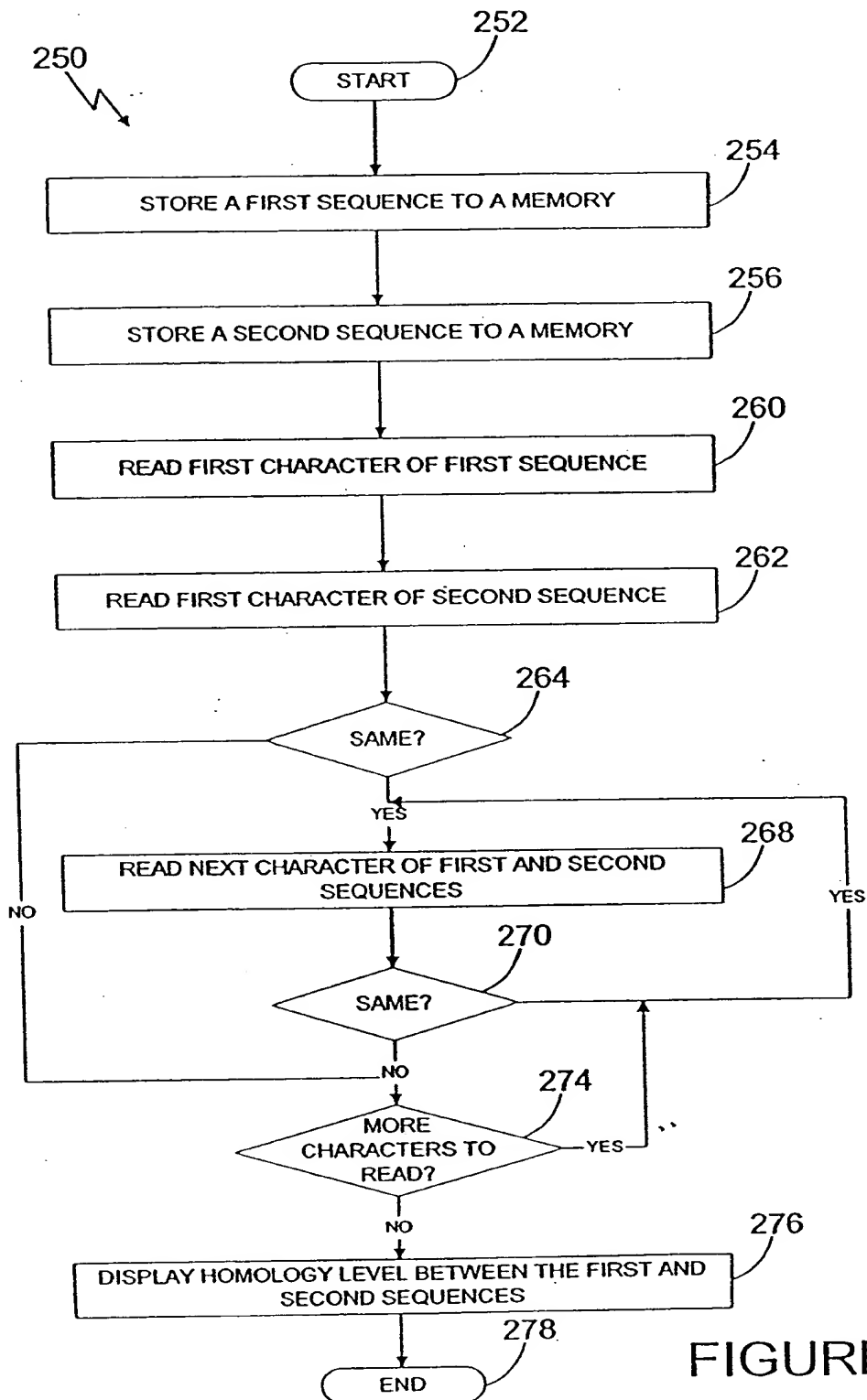


FIGURE 3

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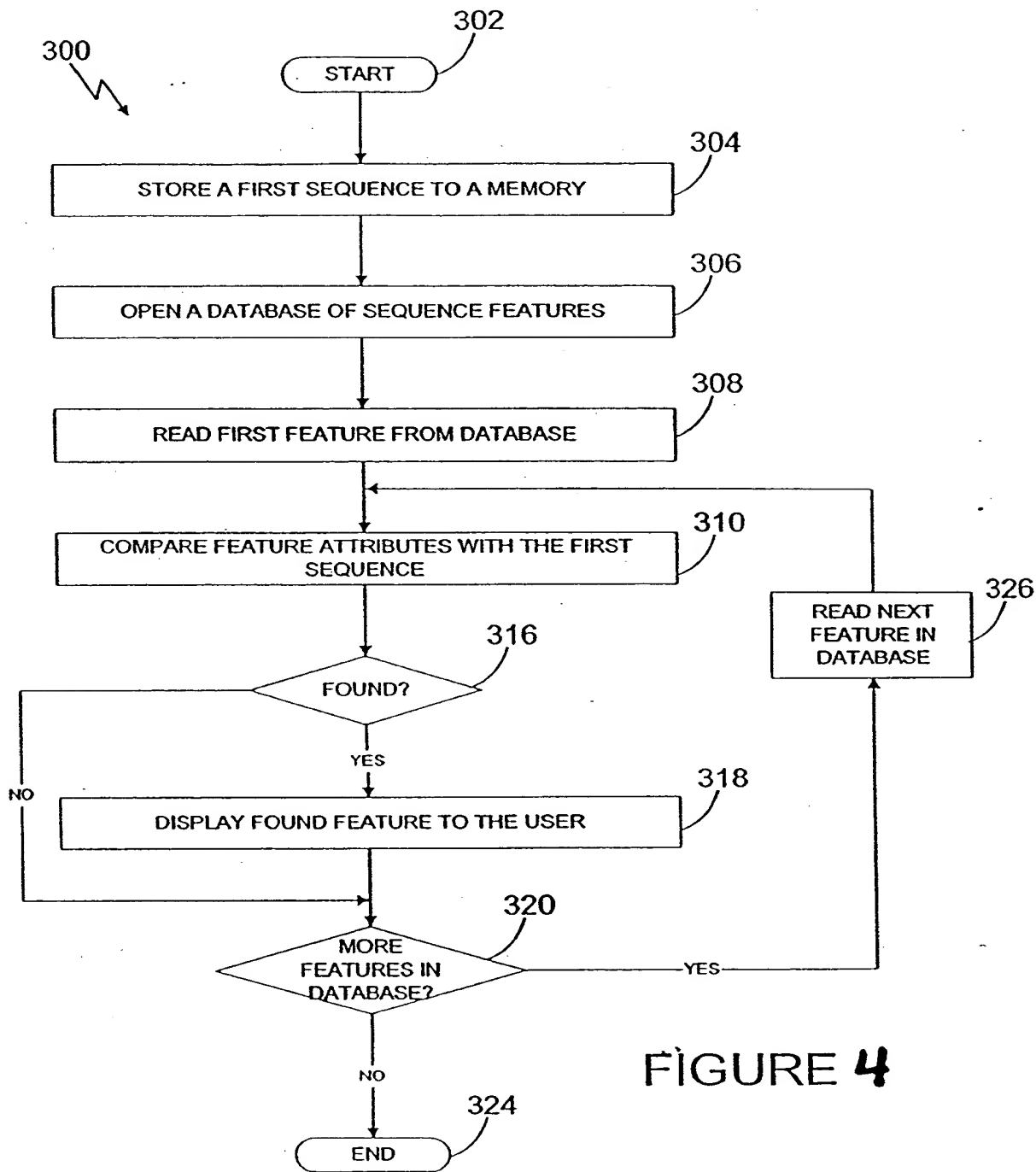


FIGURE 4

**Thermococcus GUSL5 Amidase**

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1  ATG ACC GGC ATC GAA TGG AAC CAC GAG ACC TTT TCT AAG TTC GCC TAC CTG GGC GAC CCG  60
1  Met Thr Gly Ile Glu Trp Asn His Glu Thr Phe Ser Lys Phe Ala Tyr Leu Gly Asp Pro  20

61  AGG ATA CGG GGA AAC TTA ATC GCG TAC ACC CTG ACG AAG GCC AAC ATG AAG GAC AAC AAG
120

21  Arg Ile Arg Gly Asn Leu Ile Ala Tyr Thr Leu Thr Lys Ala Asn Met Lys Asp Asn Lys  40

121  TAC GAG AGC ACG GTT GTT GAA GAC CTT GAA ACG GGC TCA AGG CGC TTC ATC GAG AAC  180
41  Tyr Glu Ser Thr Val Val Glu Asp Leu Glu Thr Gly Ser Arg Phe Ile Glu Asn  60

181  GCC TCA ATG CCG AGG ATT TCG CCA GAC GGC AGA AAG CTC GCC TTC ACC TGC TTT AAC GAG  240
61  Ala Ser Met Pro Arg Ile Ser Pro Asp Gly Arg Lys Leu Ala Phe Thr Cys Phe Asn Glu  80

241  GAG AAG AAG GAG ACC GAG ATA TGG GTG GCC GAT ATC CAG ACC CTG AGC GCC AAG AAA GTC
300

81  Glu Lys Lys Glu Thr Glu Ile Trp Val Ala Asp Ile Gln Thr Leu Ser Ala Lys Lys Val  100

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**FIG. 5A**

301 CTC TCA ACT AAA AAC GTC CGC TCG ATG CAG TGG AAC GAC GAT TCA AGG AGA CTC TTA GTT 360  
 101 Leu Ser Thr Lys Asn Val Arg Ser Met Gln Trp Asn Asp Asp Ser Arg Arg Leu Leu Val 120  
  
 361 GTC GGC TTC AAG AGG AGG GAC GAT GAG GAC TTC GTC TTT GAC GAC GAC GTC CCG GTC TGG 420  
 121 Val Gly Phe Lys Arg Arg Asp Asp Glu Asp Phe Val Phe Asp Asp Val Pro Val Trp 140  
  
 421 TTC GAC AAT ATG GGA TTC TTT GAT GGA GAG AAG ACG ACG TTC TGG GTT CTT GAC ACT GAG 480  
 141 Phe Asp Asn Met Gly Phe Phe Asp Gly Glu Lys Thr Thr Phe Trp Val Leu Asp Thr Glu 160  
  
 481 GCC GAG GAG ATA ATC GAG CAG TTC GAG AAG CCG AGG TTT TCG AGT GGC CTC TGG CAC GGC 540  
 161 Ala Glu Glu Ile Ile Glu Gln Phe Glu Lys Pro Arg Phe Ser Ser Gly Leu Trp His Gly 180  
  
 541 GAT GCG ATA GTT GTG AAC GTC CCG CAC CGC GAG GGG AGC AAG CCT GCC CTG TTC AAG TTC 600  
 181 Asp Ala Ile Val Val Asn Val Pro His Arg Glu Gly Ser Lys Pro Ala Leu Phe Lys Phe 200  
  
 601 TAC GAC ATA GTC CTA TGG AAG GAC GGG GAG GAA GAG AAG CTC TTC GAG AGG GTC TCC TTC 660  
 201 Tyr Asp Ile Val Leu Trp Lys Asp Gly Glu Glu Lys Lys Leu Phe Glu Arg Val Ser Phe 220  
  
 661 GAG GCG GTT GAC TCC GAC GGA AAG AGA ATA CTC CTG AGG GGC AAG AAA AAA AAG CCG TTC  
 720  
 221 Glu Ala Val Asp Ser Asp Gly Lys Arg Ile Leu Leu Arg Gly Lys Lys Lys Arg Phe 240

FIG. 5B

# TABLE 1

721	ATC AGC GAG CAC GAC TGG CTG TAC CTC TGG GAC GGC GAG CTT AAA CCG ATC TAC GAG GGC	780
241	Ile Ser Glu His Asp Trp Leu Tyr Leu Trp Asp Gly Glu Leu Lys Pro Ile Tyr Glu Gly	260
781	CCG CTC GAC GTC TGG GAA GCC AAG CTC ACG GAA GGA AAG GTC TAC TTC CTC ACT CCA GAT	840
261	Pro Leu Asp Val Trp Glu Ala Lys Leu Thr Glu Gly Lys Val Tyr Phe Leu Thr Pro Asp	280
841	GGC GGC AGG GTA AAC CTC TGG CTC TGG GAC GGG AAG GCC GAG CGT GTT ACC GGC GAC	900
281	Ala Gly Arg Val Asn Leu Trp Leu Trp Asp Gly Lys Ala Glu Arg Val Val Thr Gly Asp	300
901	CAC TGG ATT TAC GGG CTT GAC GTC AGC GAT GGC AAA GCA TTG CTC CTC ATC ATG ACC GCC	960
301	His Trp Ile Tyr Gly Leu Asp Val Ser Asp Gly Lys Ala Leu Leu Ile Met Thr Ala	320
961	ACG AGG ATA GGC GAG CTC TAC CTC TAC GAC GGC GAG CTG AAA CAG GTC ACC GAA TAC AAC	1020
321	Thr Arg Ile Gly Glu Leu Tyr Leu Tyr Asp Gly Glu Leu Lys Gln Val Thr Glu Tyr Asn	340
1021	GGG CCG ATA TTC AGG AAG CTC AAG ACC TTC GAG CCG AGG CAC TTC CGC TTC AAG AGC AAA	1080
341	Gly Pro Ile Phe Arg Lys Leu Lys Thr Phe Glu Pro Arg His Phe Arg Phe Lys Ser Lys	360
1081	GAC CTC GAG ATA GAC GGC TGG TAC CTC AGG CCG GAG GTT AAA GAG GAG AAG GCC CCG GTG	1140
361	Asp Leu Glu Ile Asp Gly Trp Tyr Leu Arg Pro Glu Val Lys Glu Glu Lys Ala Pro Val	380

FIG. 5C

1141 ATA GTC TTC GTC CAC GGC GGG CCG AAG GGC ATG TAC GGA CAC CGC TTC GTC TAC GAG ATG

1200

381 Ile Val Phe Val His Gly Gly Pro Lys Gly Met Tyr Gly His Arg Phe Val Tyr Glu Met 400

1201 CAG CTG ATG GCG AGC AAG GGC TAC TAC GTC GTC TTC GTG AAC CCG CGC GGC AGC GAC GGC

1260

401 Gln Leu Met Ala Ser Lys Gly Tyr Tyr Val Phe Val Asn Pro Arg Gly Ser Asp Gly 420

1261 TAT AGC GAA GAC TTC GCG CTC CGC GTC CTG GAG AGG ACT GGC TTG GAG GAC TTT GAG GAC

1320

421 Tyr Ser Glu Asp Phe Ala Leu Arg Val Leu Glu Arg Thr Gly Leu Glu Asp Phe Glu Asp 440

1321 ATA ATG AAC GGC ATC GAG GAG TTC TTC AAG CTC GAA CCG CAG GCC GAC AGG GAG CGC GTT

1380

441 Ile Met Asn Gly Ile Glu Glu Phe Phe Lys Leu Glu Pro Gln Ala Asp Arg Glu Arg Val 460

1381 GGA ATA ACG GGC ATA AGC TAC GGC GGC TTC ATG ACC AAC TGG GCC TTG ACT CAG AGC GAC

1440

461 Gly Ile Thr Gly Ile Ser Tyr Gly Gly Phe Met Thr Asn Trp Ala Leu Thr Gln Ser Asp 480

1441 CTC TTC AAG GCA GGA ATA AGC GAG AAC GGC ATA AGC TAC TGG CTC ACC AGC TAC GCC TTC

1500

481 Leu Phe Lys Ala Gly Ile Ser Glu Asn Gly Ile Ser Tyr Trp Leu Thr Ser Tyr Ala Phe 500

FIG. 5D



1501 TCG GAC ATA GGG CTC TGG TAC GAC GTC GAG GTC ATC GGG CCA AAT CCG TTA GAG AAC GAG  
1560

501 Ser Asp Ile Gly Leu Trp Tyr Asp Val Glu Val Ile Gly Pro Asn Pro Leu Glu Asn Glu 520

1561 AAC TTC AGG AAG CTC AGC CCG CTG TTC TAC GCT CAG AAC GTG AAG GCG CCG ATA CTC CTA  
1620

521 Asn Phe Arg Lys Leu Ser Pro Leu Phe Tyr Ala Gln Asn Val Lys Ala Pro Ile Leu Leu 540

1621 ATC CAC TCG CTT GAG GAC TAC CGC TGT CCG CTC GAC CAG AGC CTT ATG TTC TAC AAC GTG  
1680

541 Ile His Ser Leu Glu Asp Tyr Arg Cys Pro Leu Asp Gln Ser Leu Met Phe Tyr Asn Val 560

1681 CTC AAG GAC ATG GGC AAG GAA GCC TAC ATA GCG ATA TTC AAG CGC GGC GCC CAC GGC CAC  
1740

561 Leu Lys Asp Met Gly Lys Glu Ala Tyr Ile Ala Ile Phe Lys Arg Gly Ala His Gly His 580

1741 AGC GTC CGC GGA AGC CCG AGG CAC AGG CCG AAG CGC TAC AGG CTC TTC ATA GAG TTC TTC  
1800

581 Ser Val Arg Gly Ser Pro Arg His Arg Pro Lys Arg Tyr Arg Leu Phe Ile Glu Phe Phe 600

1801 GAG CGC AAG CTC AAG AAG TAC GAG GAG GGC TTT GAG GTA GAG AAG ATA CTC AAG GCG AAT  
1860

601 Glu Arg Lys Leu Lys Tyr Glu Glu Gly Phe Glu Val Glu Lys Ile Leu Lys Gly Asn 620

1861 GGG AAC TGA 1869

621 Gly Asn End 623

FIG. 5E

T07260" E0399660

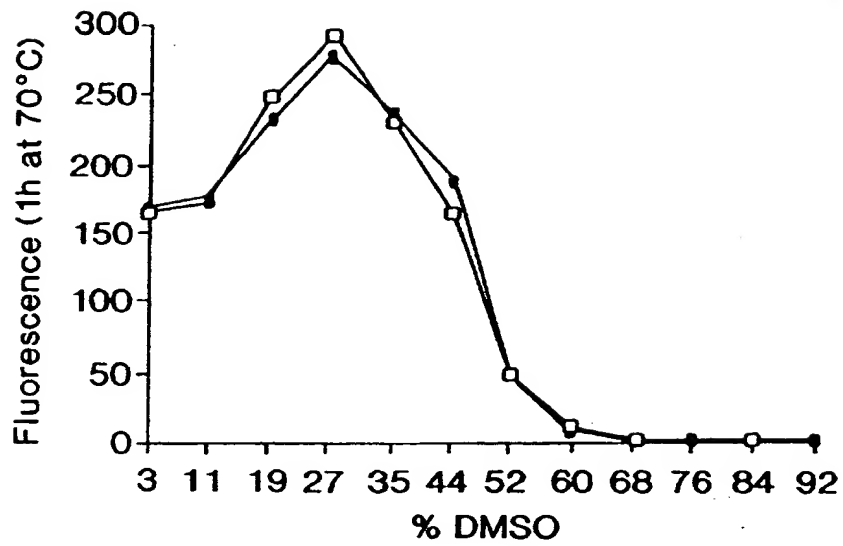


FIG. 6

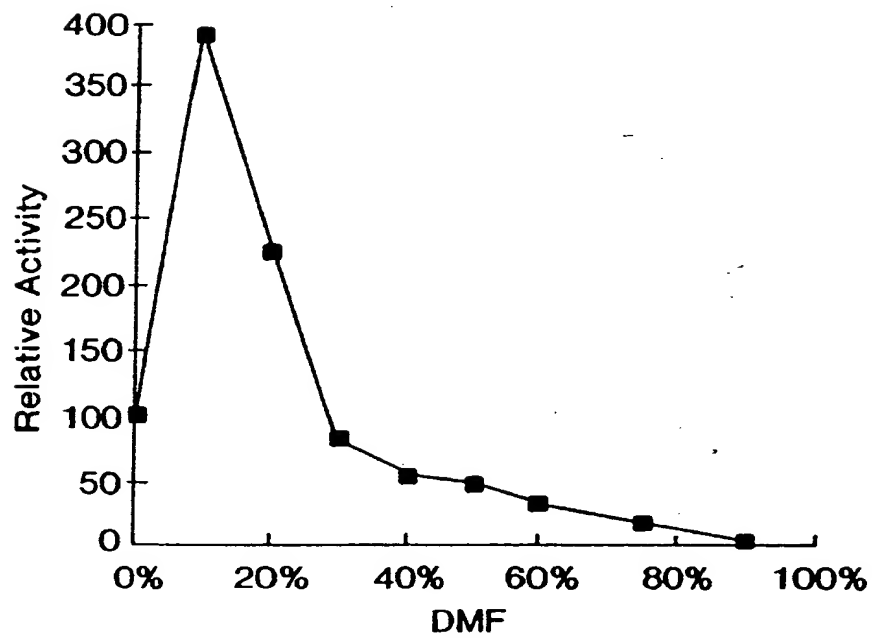


FIG. 7